

## SEQUENCE LISTING

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<120> ICAM-4 BINDING SITES

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<160> 40

<170> PatentIn version 3.3

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<211> 241

<212> PRT

<213> Homo sapiens

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Pro	Phe	Trp	Val	Arg	Met	Ser	Pro	Glu	Phe	Val	Ala	Val	Gln	Pro	Gly
			20					25					30		

Lys	Ser	Val	Gln	Leu	Asn	Cys	Ser	Asn	Ser	Cys	Pro	Gln	Pro	Gln	Asn
			35				40					45			

Ser	Ser	Leu	Arg	Thr	Pro	Leu	Arg	Gln	Gly	Lys	Thr	Leu	Arg	Gly	Pro
			50				55				60				

Gly	Trp	Val	Ser	Tyr	Gln	Leu	Leu	Asp	Val	Arg	Ala	Trp	Ser	Ser	Leu
65					70					75					80

Ala	His	Cys	Leu	Val	Thr	Cys	Ala	Gly	Lys	Thr	Arg	Trp	Ala	Thr	Ser
				85					90					95	

Arg	Ile	Thr	Ala	Tyr	Lys	Pro	Pro	His	Ser	Val	Ile	Leu	Glu	Pro	Pro
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Val	Leu	Lys	Gly	Arg	Lys	Tyr	Thr	Leu	Arg	Cys	His	Val	Thr	Gln	Val
			115					120					125		

Phe Pro Val Gly Tyr Leu Val Val Thr Leu Arg His Gly Ser Arg Val  
 130 135 140  
 Ile Tyr Ser Glu Ser Leu Glu Arg Phe Thr Gly Leu Asp Leu Ala Asn  
 145 150 155 160  
 Val Thr Leu Thr Tyr Glu Phe Ala Ala Gly Pro Arg Asp Phe Trp Gln  
 165 170 175  
 Pro Val Ile Cys His Ala Arg Leu Asn Leu Asp Gly Leu Val Val Arg  
 180 185 190  
 Asn Ser Ser Ala Pro Ile Thr Leu Met Leu Ala Trp Ser Pro Ala Pro  
 195 200 205  
 Thr Ala Leu Ala Ser Gly Ser Ile Ala Ala Leu Val Gly Ile Leu Leu  
 210 215 220  
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<400> 9  
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<210> 10  
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<213> Homo sapiens

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gcgcactgcc	tcgtgacctg	cgcaggaaaa	acacgctggg	ccacctccag	gatcaccgcc	300
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atgctcgctt	ggagccccgc	gcccacagct	ttggcctccg	gttccatcgc	tgcccttgta	660
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<400> 16  
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<210> 19  
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<400> 24  
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<400> 26  
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<400> 39  
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ccc ttc tgg gtg cgc atg agc ccg gag ttc gtg gct gtg cag ccg ggg	96
Pro Phe Trp Val Arg Met Ser Pro Glu Phe Val Ala Val Gln Pro Gly	
20 25 30	
aag tca gtg cag ctc aat tgc agc aac agc tgt ccc cag ccg cag aat	144
Lys Ser Val Gln Leu Asn Cys Ser Asn Ser Cys Pro Gln Pro Gln Asn	
35 40 45	
tcc agc ctc cgc acc ccg ctg cgg caa ggc aag acg ctc aga ggg ccg	192
Ser Ser Leu Arg Thr Pro Leu Arg Gln Gly Lys Thr Leu Arg Gly Pro	
50 55 60	
ggt tgg gtg tct tac cag ctg ctc gac gtg agg gcc tgg agc tcc ctc	240
Gly Trp Val Ser Tyr Gln Leu Leu Asp Val Arg Ala Trp Ser Ser Leu	
65 70 75 80	
gcg cac tgc ctc gtg acc tgc gca gga aaa aca cgc tgg gcc acc tcc	288
Ala His Cys Leu Val Thr Cys Ala Gly Lys Thr Arg Trp Ala Thr Ser	
85 90 95	
agg atc acc gcc tac aaa ccg ccc cac agc gtg att ttg gag cct ccg	336
Arg Ile Thr Ala Tyr Lys Pro Pro His Ser Val Ile Leu Glu Pro Pro	
100 105 110	
gtc tta aag ggc agg aaa tac act ttg cgc tgc cac gtg acg cag gtg	384
Val Leu Lys Gly Arg Lys Tyr Thr Leu Arg Cys His Val Thr Gln Val	
115 120 125	
ttc ccg gtg ggc tac ttg gtg gtg acc ctg agg cat gga agc cgg gtc	432
Phe Pro Val Gly Tyr Leu Val Val Thr Leu Arg His Gly Ser Arg Val	
130 135 140	
atc tat tcc gaa agc ctg gag cgc ttc acc ggc ctg gat ctg gcc aac	480
Ile Tyr Ser Glu Ser Leu Glu Arg Phe Thr Gly Leu Asp Leu Ala Asn	
145 150 155 160	
gtg acc ttg acc tac gag ttt gct gct gga ccc cgc gac ttc tgg cag	528
Val Thr Leu Thr Tyr Glu Phe Ala Ala Gly Pro Arg Asp Phe Trp Gln	
165 170 175	
ccc gtg atc tgc cac gcg cgc ctc aat ctc gac ggc ctg gtg gtc cgc	576
Pro Val Ile Cys His Ala Arg Leu Asn Leu Asp Gly Leu Val Val Arg	
180 185 190	
aac agc tcg gca ccc att aca ctg atg ctc gct tgg agc ccc gcg ccc	624
Asn Ser Ser Ala Pro Ile Thr Leu Met Leu Ala Trp Ser Pro Ala Pro	
195 200 205	

